

RAW SEQUENCE LISTING

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Application Serial Number: 10/576, 093
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/576,093

DATE: 02/22/2007

TIME: 13:44:33

Input Set : E:\22339-Sequence Listing.txt
 Output Set: N:\CRF4\02222007\J576093.raw

3 <110> APPLICANT: Roche Molecular Systems
 5 <120> TITLE OF INVENTION: Method for Distinguishing MLL-PTD-Positive AML from Other
 AML
 6 Subtypes
 8 <130> FILE REFERENCE: 22339-US
 10 <140> CURRENT APPLICATION NUMBER: US 10/576,093
 C--> 11 <141> CURRENT FILING DATE: 2006-04-13
 13 <150> PRIOR APPLICATION NUMBER: PCT/EP2004/012464
 14 <151> PRIOR FILING DATE: 2003-11-04
 16 <160> NUMBER OF SEQ ID NOS: 239
 18 <170> SOFTWARE: PatentIn version 3.4
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 491
 22 <212> TYPE: DNA
 23 <213> ORGANISM: Homo sapiens
 26 <220> FEATURE:
 27 <221> NAME/KEY: misc_feature
 28 <223> OTHER INFORMATION: n = any nucleotide
 30 <400> SEQUENCE: 1
 31 gaagcccaaa aactcttgg acaatataag gaagaaaagca aaaaggctct tccaccagaa 60
 33 aagaaaacaga acactggctc aaagaaaaagc aataaaaaata agagtggcaa gaaccagttt 120
 35 aacagaggtg gtggccatag aggacgttgg ggattcaata tgcgtgggtt aaatttcaga 180
 37 ggaggagccc ctggaaatcg tggcggatataataggaggg gcaacatgcc acagagaggt 240
 39 ggtggcgttg gaggaagtgg tggaaatcggc tatccataacc ctcgtgcccc tgttttcct 300
 41 ggcgtgtt gttactcaaa cagagggaac tacaacagag gtggaaatgcc caacagaggg 360
 43 aactacaacc agaacttcag aggacgagga aacaatcgtg gctacaaaaaa tcaatctcag 420
 45 ggctacaacc agtggcagca gggtcaattc tgggtcaga agccatggag tcagcattat 480
 47 caccaaggat a 491
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 51 <211> LENGTH: 307
 52 <212> TYPE: DNA
 53 <213> ORGANISM: Homo sapiens
 55 <400> SEQUENCE: 2
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 58 cgctttctac actgttattac ataaataaaat taaataaaat aaccccgggc aagacttttc 120
 60 tttaaggat gactacagac attaataat cgaagtaatt ttgggtgggg agaagaggca 180
 62 gattcaattt tcttaacca gtctgaagg tcatttatga tacaagaa gatgaaaatg 240
 64 gaagtggcaa tataagggta tgaggaaggc atgcctggac aaacccttct tttaagatgt 300
 66 gtcttca 307
 69 <210> SEQ ID NO: 3
 70 <211> LENGTH: 519
 71 <212> TYPE: DNA
 72 <213> ORGANISM: Homo sapiens
 74 <400> SEQUENCE: 3

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75	gttggaaact ttaatagcgt tgcaacgaaa tcctatatcc agtttcctgt aatattaattg	60
77	aagaaaaata catccaaata aagactttat tattaacaga ccagatagca tcagaaatca	120
79	tgtgactgtt atgattatca gaatatgtct taactttta gggcaaagg aacactgaaa	180
81	gttctagctt aagtgtgaa acttttgtgg gaaaaaaaaa tcactttga aactcagact	240
83	tcagtgtata cccaaataatt taaaattatg tgaaatgttt taaatttgg aactcgtaat	300
85	tactgtttta atgattcagt ttcttcagag tggtaattgt ataaaaattgc tattgcagct	360
87	ttatattcaa tatgatgtgc ctgtaaacca aggagtttc cccgtttgtaaaaaagacatt	420
89	gtagataatt gaatgttga ttttagaaag gtcatttagtt tcttgttacaaattttgtta	480
91	gtctggttt tggcgttatt cgggttaat attgttctt	519
94	<210> SEQ ID NO: 4	
95	<211> LENGTH: 140	
96	<212> TYPE: DNA	
97	<213> ORGANISM: Homo sapiens	
99	<400> SEQUENCE: 4	
100	ctaccttatcc tgaatggct gtcattgtct gcctttaaaa tccttcctct ttcttcctcc	60
102	tctattctct aaataatgtat ggggctaagt tataacccaaa gtcacttta caaaatattt	120
104	cctcagtaact ttgcagaaaa	140
107	<210> SEQ ID NO: 5	
108	<211> LENGTH: 425	
109	<212> TYPE: DNA	
110	<213> ORGANISM: Homo sapiens	
112	<400> SEQUENCE: 5	
113	gtcatatcat ttcaactgtct aggctacaac aggattctag gtggaggttg tgcattgtgt	60
115	ccttttatac tgatctgtga ttaaaggcgt aatattttaa gatggactgg gaaaaacatc	120
117	aactcctgaa gttagaaata agaatggttt gtaaaaatcca cagctatatac ctgatgctgg	180
119	atggattttaa tcttggtag tcttcaactg gttatgtga aatagttctg ccacccctgaa	240
121	cgcaccactg ccaatgtctg acgtactgca tttggccctt gagccaggtg gatgtttacc	300
123	gtgtgttata taacttcctg gtccttcac tgaacatgcc tagtccaaca tttttccca	360
125	gtgagtcaca tcctgggatc cagtgtataa atccaatatac atgtcttgcataattctt	420
127	ccaaa	425
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131	<211> LENGTH: 454	
132	<212> TYPE: DNA	
133	<213> ORGANISM: Homo sapiens	
135	<400> SEQUENCE: 6	
136	caagctatgg aataccctgg gtgtgtgcaa atacactgtc caggatgaga gccactcaga	60
138	gtgggtgtct tgggtccgct tctcgcccaa cagcagcaac cctatcatcg tctccgtgg	120
140	ctgggacaag ctggcataagg tatggaacct ggctactgc aagctgaaga ccaaccacat	180
142	tggccacaca ggctatctga acacggtagc tgcattctcca gatggatccc tctgtgcttc	240
144	tggaggcaag gatggccagg ccatgtttag ggtatctcaac gaaggcaaac acctttacac	300
146	gctagatggt ggggacatca tcaacggcct gtgcattcagc cctaaaccgct actggctgtg	360
148	tgctgccaca ggccccagca tcaagatctg ggattttagag ggaaagatca ttgttagatga	420
150	actgaagcaa gaagttatca gtaccagcag caag	454
153	<210> SEQ ID NO: 7	
154	<211> LENGTH: 373	
155	<212> TYPE: DNA	
156	<213> ORGANISM: Homo sapiens	
158	<400> SEQUENCE: 7	
159	atcagggtat ttgttccacc ttggccaggc ctcctcggag aagcttgc cccgtgtggg	60

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161	aggcacggag	ccggacttgg	catggtaact	cagtaccgc	tgcagtgtcg	ccatgactga	120
163	tcatggctct	tgcattttt	ggtaaatgg	gactccgg	tcctgtcagg	gtgtccccca	180
165	tgcctggaaag	aggagcttgg	ggctgccagc	cctggcgccg	gcacagcctg	ggcctccct	240
167	tccctcaaggc	cagggttctt	cctcctgtcg	tgggttcatt	tgccaggctc	agggcaggtc	300
169	tggacagctg	tgactcttct	caagccagga	ctaccgacca	gccggctatg	ggcacattac	360
171	gtgaccactg	gcc					373
174	<210>	SEQ ID NO:	8				
175	<211>	LENGTH:	494				
176	<212>	TYPE:	DNA				
177	<213>	ORGANISM:	Homo sapiens				
179	<400>	SEQUENCE:	8				
180	agtgcggaca	ggacgggtcat	tgattacaac	ggggaaacgca	cgctggatgg	tttaagaaaa	60
182	tccctagaga	gcgggtggca	agatggggca	ggggatgtt	acgacccctcg	ggacctcgaa	120
184	gaagcagagg	agccagacat	ggaggaagac	gatgaccaga	aagctgtgaa	agatgaactg	180
186	taataacgcaa	agccggaccc	gggcgctg	gagaccctc	gggggctgca	cacccagcag	240
188	cagcgcacgc	ctccgaagcc	tgccgcctcg	cttgaaggag	ggcgtcgccg	gaaacccaag	300
190	gaacctctct	gaagtacac	ctcaccccta	cacaccgtcc	gttcacccccc	gtctcttcct	360
192	tctgtttt	ggtttttgg	aaacccggat	cctactctag	gcagccccc	ttggtgggct	420
194	tgtttctgaa	accatgtat	tacttttca	tacatgagtc	tgtccagagt	gttgctacc	480
196	gtgttcggag	tctc					494
199	<210>	SEQ ID NO:	9				
200	<211>	LENGTH:	434				
201	<212>	TYPE:	DNA				
202	<213>	ORGANISM:	Homo sapiens				
204	<400>	SEQUENCE:	9				
205	cctacttcgg	tatctatgac	actgcaaagg	aatgcttcc	ggatcccaag	aacactcaca	60
207	tgcgtcatcg	ctggatgtat	gcacagactg	tcactgtgt	tgccgggttg	acttcctatc	120
209	catttgacac	cgttcgccgc	cgcatgtat	tgcagtccgg	gcgcaaaagg	actgacatca	180
211	tgtacacagg	cacgttgcac	tgctggcgga	agattgtcg	tgtatggaa	ggcaaaagctt	240
213	ttttcaagg	tgcatggtcc	aatgttctca	gaggcatgg	ttgtgtttt	gtgttgtct	300
215	tgtatgtat	aatcaagaag	tacacataag	ttatcccta	ggattttcc	ccctgtgaac	360
217	aggcatgtt	tattctataa	cacaatctt	agcattctt	acagactcct	ggctgtcagt	420
219	ttctcagtgg	caac					434
222	<210>	SEQ ID NO:	10				
223	<211>	LENGTH:	416				
224	<212>	TYPE:	DNA				
225	<213>	ORGANISM:	Homo sapiens				
227	<400>	SEQUENCE:	10				
228	gttggttcaa	acttttgg	gcacggactg	tcaatgttct	ggaaagtgg	cagcgcattcc	60
230	tgcagggtt	ctcccttct	gtcttttgg	gaaccagg	tcttcagg	ggctctagg	120
232	actgcccggc	tgtttcagcc	aggaaaggcc	aaatcaagag	ttagatgtat	aaatgttata	180
234	aatagaaaaaa	gtggagttgg	tgaatcggtt	gttcttcct	cacatttgg	tgattgtat	240
236	aagggttttta	gcatgttcc	ccttttctt	accctccct	ttgttcttct	attaatcaag	300
238	agaaacttca	aagttaatgg	gttgcgg	tctcacaggc	tgagaactcg	ttcacctcca	360
240	agcatttcat	aaaaaagctg	tttcttataa	atcataaaaa	ctctcaccat	gtatgt	416
243	<210>	SEQ ID NO:	11				
244	<211>	LENGTH:	415				
245	<212>	TYPE:	DNA				
246	<213>	ORGANISM:	Homo sapiens				

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248 <400> SEQUENCE: 11		
249 accccctgtg aagcccaaga tcgtcaaaaaa gagaaccaag aagttcatcc ggcaccagtc	60	
251 agaccgatat gtcaaaatta agcgtaactg gcggaaaccc agaggcattg acaacagggt	120	
253 tcgtagaaga ttcaagggcc agatcttcat gccaacatt gttatggaa gcaacaaaaa	180	
255 aacaaagcac atgctgccc gtggcttccg gaagttcctg gtccacaacg tcaaggagct	240	
257 ggaagtgtcg ctgtatgtca acaaatactt ctgtgccgag atcgctcaca atgtttcctc	300	
259 caagaaccgc aaaqccatcg tggaaagagc tgcccaactg gccatcagag tcaccaaccc	360	
261 caatgccagg ctgcgcagtg aaaaaatgta gtaggcagct catgtgcacg ttttc	415	
264 <210> SEQ ID NO: 12		
265 <211> LENGTH: 423		
266 <212> TYPE: DNA		
267 <213> ORGANISM: Homo sapiens		
269 <400> SEQUENCE: 12		
270 aataatttat tcccacatct acatcagtga aagctatcta cctatcctga gtctatctta	60	
272 aaggaaaaaa aaaaaaaaaac cttatctttt gcccattttt tgaattttcc actctttcat	120	
274 taatttgttt taagctcctg ttggaaaaaa agggtagtg cattttaaat tgaccttcat	180	
276 acgctttaa aataagacaa atctacttga taatgtacct ttatgtatc tcaagttgt	240	
278 taaaaccaat aaatttgtgt tactgcagta gtaatcttgc acacacggtg atttcatgtt	300	
280 atatatgcac aqtaggcaac tgttttctta gttacagaag tttcaagctt cactttgtg	360	
282 cagtagaaac aaaagtaggc tacagtctgt gccatgttgc tgcacatgtt ctgaaaattgt	420	
284 ttt	423	
287 <210> SEQ ID NO: 13		
288 <211> LENGTH: 358		
289 <212> TYPE: DNA		
290 <213> ORGANISM: Homo sapiens		
292 <400> SEQUENCE: 13		
293 tgcttctggac cacctggac caggtcttg tctgggttgg aaaggattct caagaagaag	60	
295 aaaagacaga agccttgact tctgctaagc ggtacatcga gacggacca gccaatcgaa	120	
297 atcggcggac gcccatttacc gttgtaaagc aaggcttgc gcctccctcc ttgtgggt	180	
299 gttccttgg ctggatgtt gattactgtt ctgtggaccc cttggacagg gcatggctg	240	
301 agctggctgc ctggaggaggg gcaggggccca cccatgtcac cggtcagtgcc ttgtggaaac	300	
303 tgccttccc tcaaagaggc cttagagcga gcagagcgc tctgtatgc gtgtgtgt	358	
306 <210> SEQ ID NO: 14		
307 <211> LENGTH: 529		
308 <212> TYPE: DNA		
309 <213> ORGANISM: Homo sapiens		
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314 gggaaaagaag ctcagttgc cagcataagt gccaagaccc atcaccagca tctgtccttc	120	
316 aggtgctcg gacagaattc ttaccacagc aaagcataa gatgcttgc acgaaaatc	180	
318 agaaaacttaa ctctttgtt gcagatgtc atcagtggtt ctgtaaaaac gcagaggaaa	240	
320 agagccagaa ggtttctgtt taatgcattt tgccttatct tttttatata ctgtgtacaa	300	
322 agatttttt acacaaagaa acttaatgtt gtattaataa attcagtggtt tagttcaat	360	
324 tggatagtt cccaaaagtga agatttgtt aggaataagt gcaaattttt tttttatata	420	
326 aaaaaattct ttgaaactct taagtcttg tgcctgcaat gaaattgtac tccttgacag	480	
328 ttgatagatt atgtattctt ccatccctca aactgcatt ccactatata	529	
331 <210> SEQ ID NO: 15		
332 <211> LENGTH: 529		
333 <212> TYPE: DNA		

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334 <213> ORGANISM: Homo sapiens
 336 <400> SEQUENCE: 15

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339	ggctgcccga	atttgc	ataagtacat	ggtaaaaagt	tgtggcaaag	atggcttcca	120
341	tatccgggtg	cggctccacc	ccttccacgt	catccgcac	aacaagatgt	tgtcctgtgc	180
343	tggggctgac	aggctccaaa	caggcatgcg	aggtgcctt	ggaaagcccc	aggcactgt	240
345	ggccagggtt	cacattggcc	aagtatcat	gtccatccgc	accaagotgc	agaacaagga	300
347	catgtgatt	gaggccctgc	gcaggccaa	gttcaagttt	cctggccccc	agaagatcca	360
349	catctcaaag	aagtggggct	tcaccaagtt	caatgtgtat	gaatttgaag	acatgggtgg	420
351	tgaaaagcg	ctcatcccag	atggctgtgg	ggtcaagtac	atccccagtc	gtggccctct	480
353	ggacaagtgg	cggccctgc	actcatgagg	gttccaatg	tgctgcccc		529
356	<210>	SEQ ID NO: 16					
357	<211>	LENGTH: 393					
358	<212>	TYPE: DNA					
359	<213>	ORGANISM: Homo sapiens					
361	<400>	SEQUENCE: 16					
362	aacactcttg	tggtaaagaa	atctgatgtg	gaggcaatct	tttcgaagta	tggcaaaatt	60
364	gtgggctgt	ctgttccat	gggctttgcc	ttcggtcagt	atgttaatga	gagaaatgcc	120
365	ggggctgt	tagcaggaga	ggatggcaga	atgatgtgt	gccagggttt	agatattaa	180
368	ctggctgcag	agccaaaagt	aaaccgagga	aaagcagggtg	tgaaacgatc	tgcagcggag	240
370	atgtacggct	cctctttga	cttggactat	gacttcaac	gggactattt	tgtataggatg	300
372	tacagttacc	cagcacgtgt	acccctcct	cctcctattt	ctcggtctgt	agtgcctcgt	360
374	aaacgtcagc	gtgtatcagg	aaacacttca	cga			393
377	<210>	SEQ ID NO: 17					
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380	<213>	ORGANISM: Homo sapiens					
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385	gccaggtata	atgttaacttc	accccagcct	ttgtactaag	ctcttgcata	tggatataact	120
387	cttttaagtt	tagcccaat	atagggtat	ggaaatttcc	tgcctctgg	gttccccatt	180
389	tttactat	agaagaccag	tgataatttta	ataatgccac	caactctggc	ttagttaagt	240
391	gagagtgtg	actgtgtggc	aagagaggct	cacacccac	taggtgcaga	gagcccaggc	300
393	tttatgtttaa	aatcatgcac	ttgaaaagca	aacccat	tgcaaagaca	gcagcaagca	360
395	ttatacggc	atcttgaatg	atcccttga	aattttttt	ttgtttgtt	gtttaatca	420
397	agcctgaggc	tggtaacag	tagctacaca	cccatattgt	gtgttctgt	aatgttagct	480
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403	<211>	LENGTH: 534					
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405	<213>	ORGANISM: Homo sapiens					
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408	aggtgcagaa	tctggcg	gagcttgg	tctctgggt	ggacagcgcc	atgtccctga	60
410	tccaggcagc	caagaactt	atgaatgt	tggcagac	agtgaaggca	tcctacgtcg	120
412	cctctaccaa	atacaaaag	tcacaggta	tggcttccct	caacccat	gctgtgtcat	180
414	gaaatgtgaa	ggcaccagag	aaaaaggccat	tggtaagag	agagaaaacag	gttgagacac	240
416	agaccaagat	taaacggca	tctcagaaga	agcacgtaa	cccggtgcag	gcctcagcg	300
418	agttcaaagc	tatggacagc	atctaagtct	gcccaggccg	gcccggccca	cccctcg	360
420	ctcctgaaata	tcagtcact	aaatgaattt	gctaaataca	acactgatac		420

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:26; N Pos. 174,389
Seq#:32; N Pos. 36
Seq#:33; N Pos. 105,112,127,141
Seq#:40; N Pos. 35
Seq#:55; N Pos. 34,104,141,152,174,177,197,202,204,206,232,235,244,271,303
Seq#:55; N Pos. 312,315,349,354,384,389,400,407,415,423,499
Seq#:72; N Pos. 37,177
Seq#:86; N Pos. 297,298
Seq#:96; N Pos. 321,322
Seq#:122; N Pos. 231
Seq#:124; N Pos. 287
Seq#:156; N Pos. 277
Seq#:160; N Pos. 76
Seq#:163; N Pos. 66
Seq#:165; N Pos. 190,191,192,194,195,196,197,198,199,200,308
Seq#:191; N Pos. 182,475,514,515
Seq#:209; N Pos. 148
Seq#:216; N Pos. 57,58,59,183,303,305,316,389,404

VERIFICATION SUMMARY
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L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:606 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:120
L:614 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:360
L:744 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:775 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:60
L:777 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:120
L:936 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:0
L:1295 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:0
L:1297 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:60
L:1299 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:120
L:1301 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:180
L:1303 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:240
L:1305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:300
L:1307 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:360
L:1309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:420
L:1311 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:480
L:1684 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72 after pos.:0
L:1688 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72 after pos.:120
L:2002 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:240
L:2240 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:96 after pos.:300
L:2836 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:122 after pos.:180
L:2894 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:124 after pos.:240
L:3644 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:156 after pos.:240
L:3744 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:160 after pos.:60
L:3825 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:163 after pos.:60
L:3881 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:165 after pos.:180
L:3885 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:165 after pos.:300
L:4503 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:191 after pos.:180
L:4511 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:191 after pos.:420
L:4513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:191 after pos.:480
L:4947 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:209 after pos.:120
L:5124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:216 after pos.:0
L:5130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:216 after pos.:180
L:5134 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:216 after pos.:300
L:5136 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:216 after pos.:360